

Class 22: Microarrays (1)

Held: Tuesday, 17 November 2009

Summary: We consider our final bioinformatic technique, microarrays. Microarrays are used to study gene expression.

Related Pages:

- EBoard.
- Reading: Chapter 9.
- Due: Peer reviews of project proposals.

Notes:

- Exam 1 returned.
- Reading for Thursday: Tuggle et al. No response paper, just some prepared questions (which you should email to us).
- Two talks Thursday afternoon: Chris Tuggle on Bioinformatics and David Sloan Wilson on Trusth and Reconciliation for Group Selection.
- Everyone should have received comments on their project proposals from Sam, Vida, and some peers.
- Reminder: You will be giving quick descriptions of your projects on Thursday.

Overview:

- Microarrays.
- About Exam 1.
- Project Proposals.

Microarrays

- A technique for exploring gene expression.
- Used primarily for *comparative* gene expression.
- Clearly biotechnology.
- Why is it considered a problem in bioinformatics?

Discussion of Exam 1

- We'll quickly go over some of the more perplexing parts of exam 1.
- Feel free to visit us with questions.

Projects

- You've now received a variety of feedback on your projects.
 - Some general issues:
 - Did not explore the literature. It would be nice to see what similar things have been done (and to have confidence that you've considered whether similar things have been done).
 - Did not explore the data. There seemed to be a bit too much confidence that the data you need will be available or that certain pre-hypotheses are met.
 - We will give your groups a bit of time to revise your proposals.
 - Remember: You'll be giving quick presentations of your proposals in Thursday's class.
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